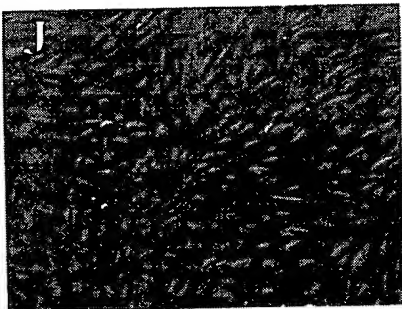
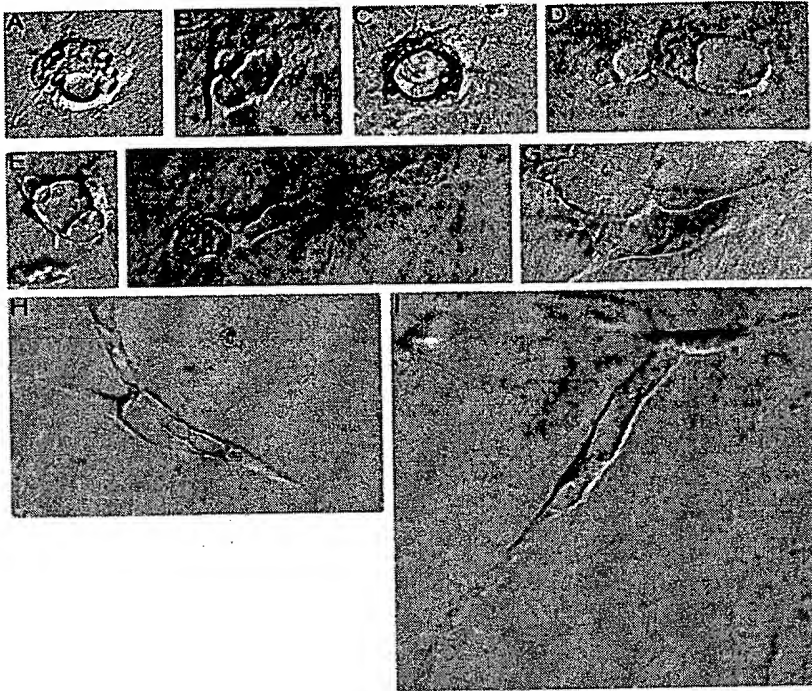


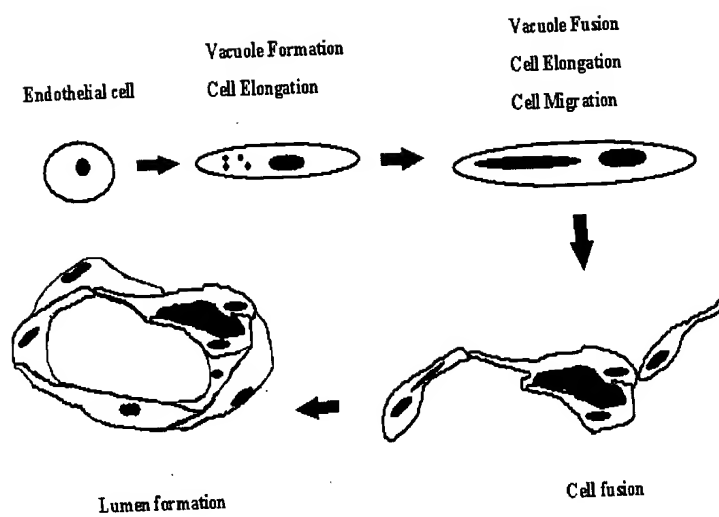
FIGURE 1



HUVEC on collagen film, 48 hr.

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FIGURE 2



Formation of three dimensional tubular networks some with interconnected lumens

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FIGURE 3: PA1 (osteonidogen)

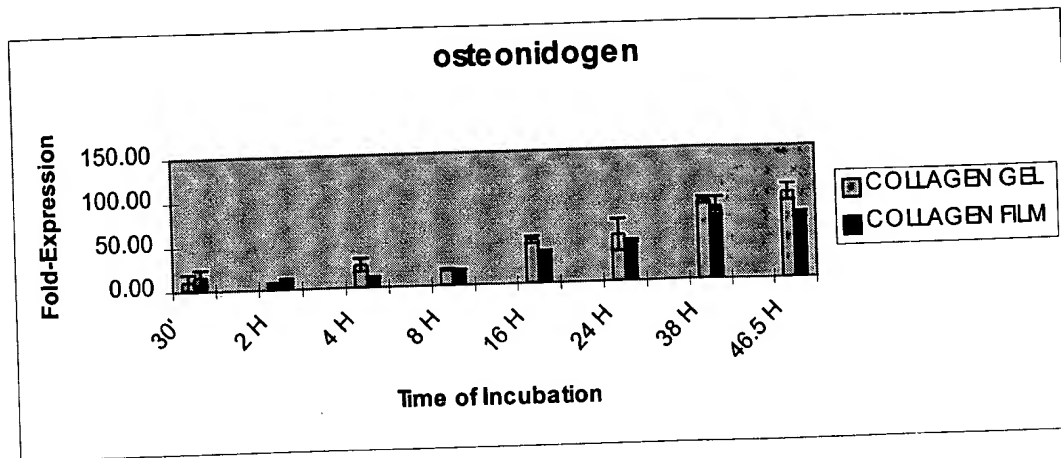
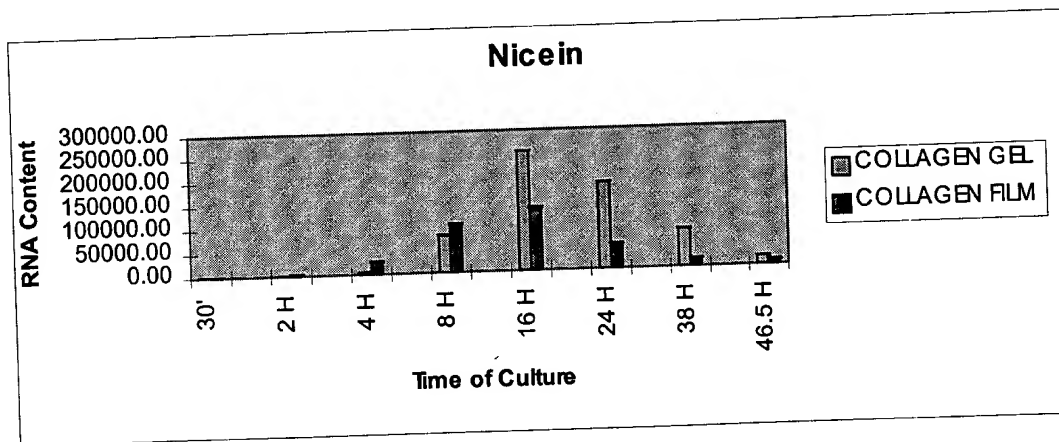


FIGURE 4: PA2 (laminin gamma-2 chain (nicein))



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FIGURE 5: PA3 (podocalyxin like protein)

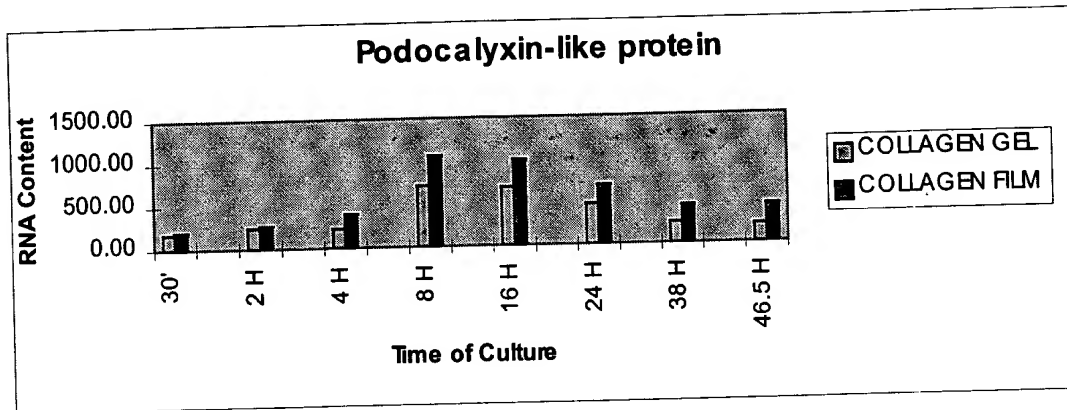


FIGURE 6: PA4 (moesin)

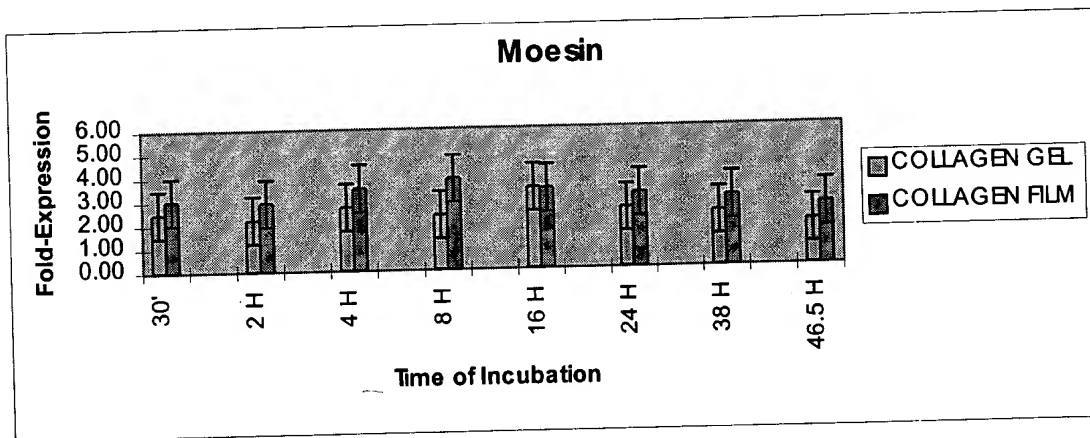


FIGURE 7: PA5 (mesoendothelial keratin K7)

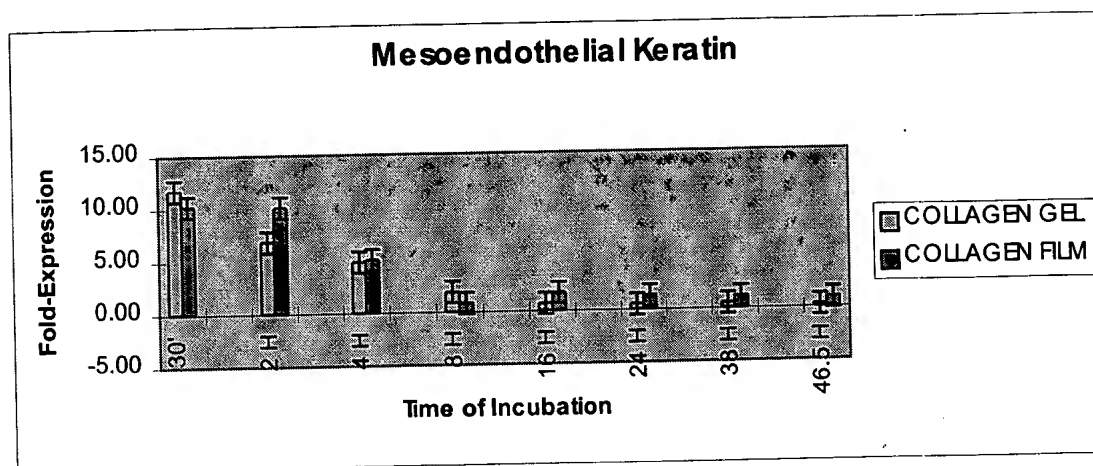
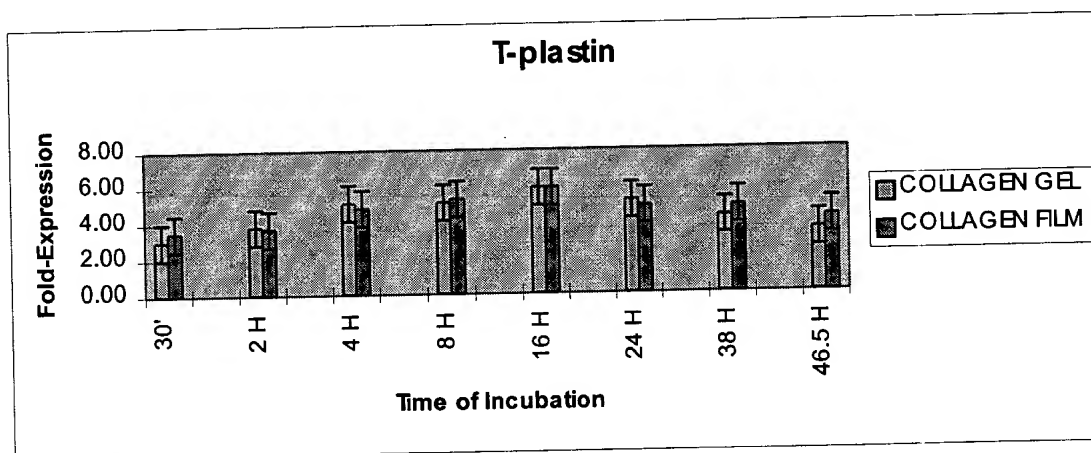


FIGURE 8: PA7 (T-plastin)



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FIGURE 9: PA8 (actin bundling protein)

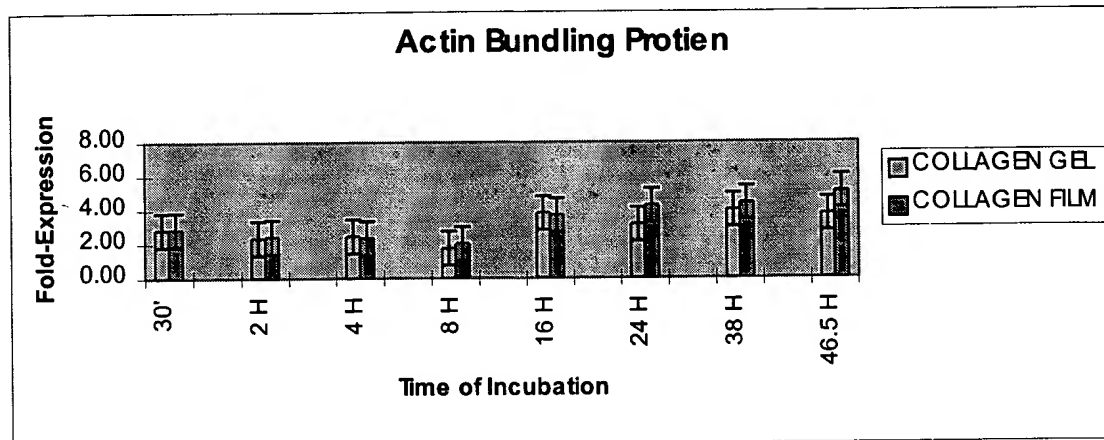


FIGURE 10: PA10 (C3VS homolog)

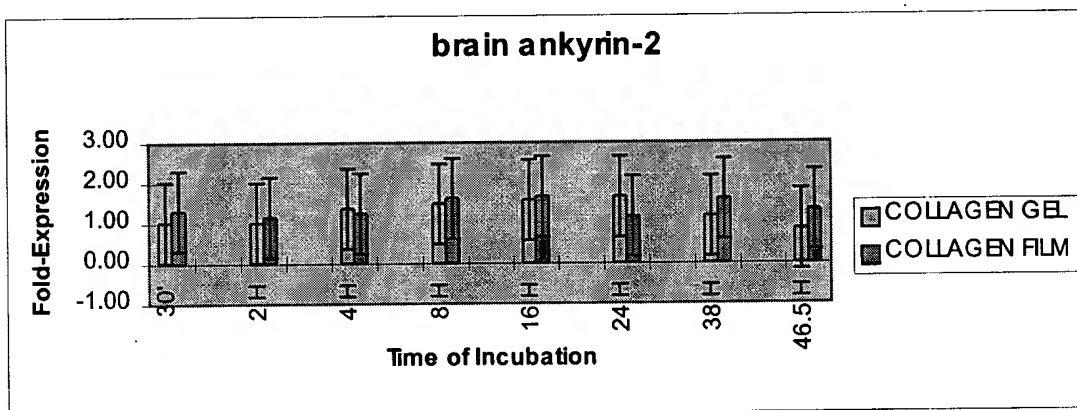


FIGURE 11: PA13 (tissue factor pathway inhibitor)

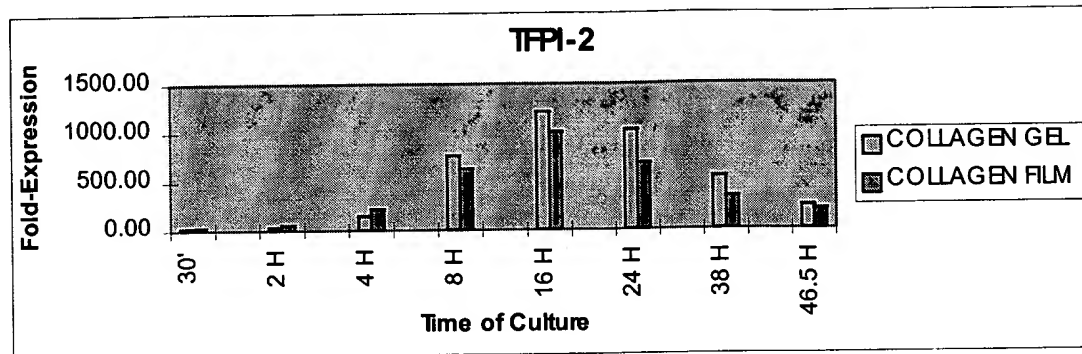


FIGURE 12: PA11 (cathepsin B)

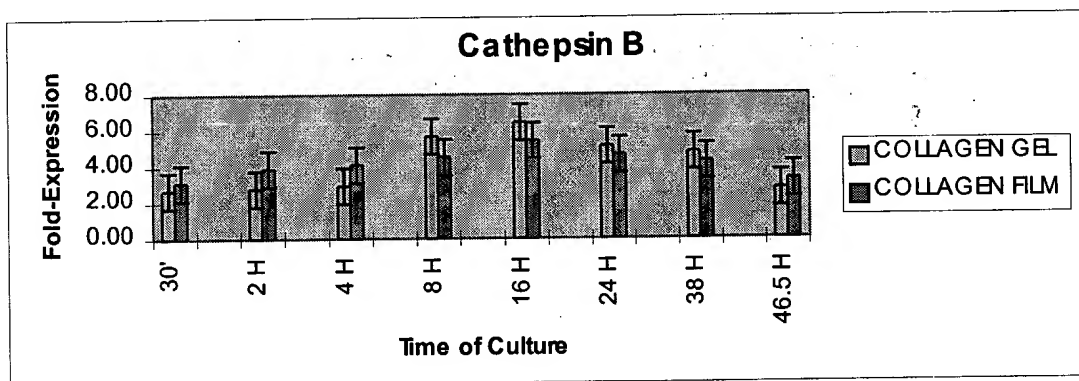


FIGURE 13: PA14 (plasminogen activator inhibitor)

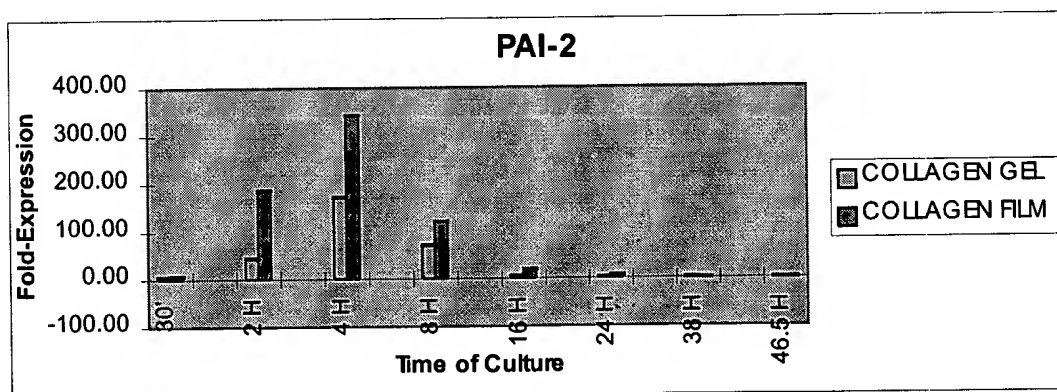


FIGURE 14: PA12 (ADAMTS-4)

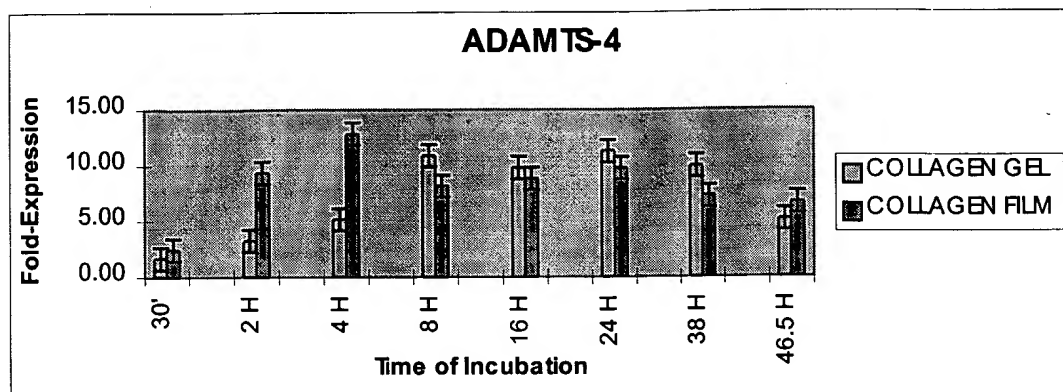
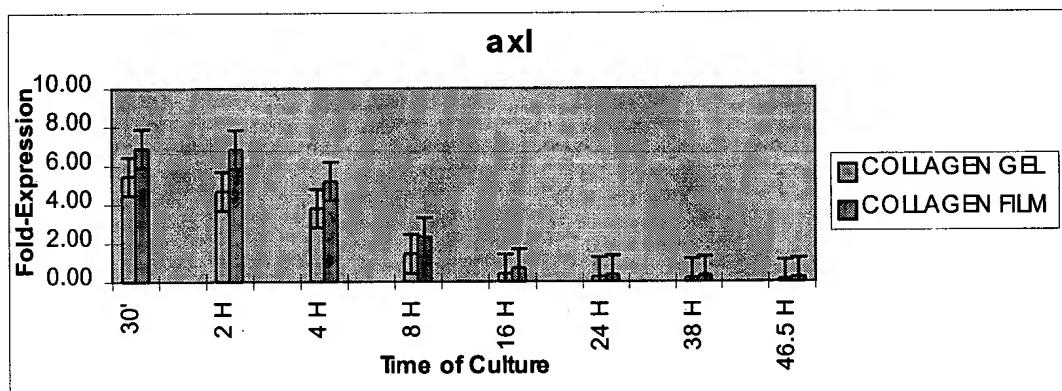


FIGURE 15: PA15 (axl)



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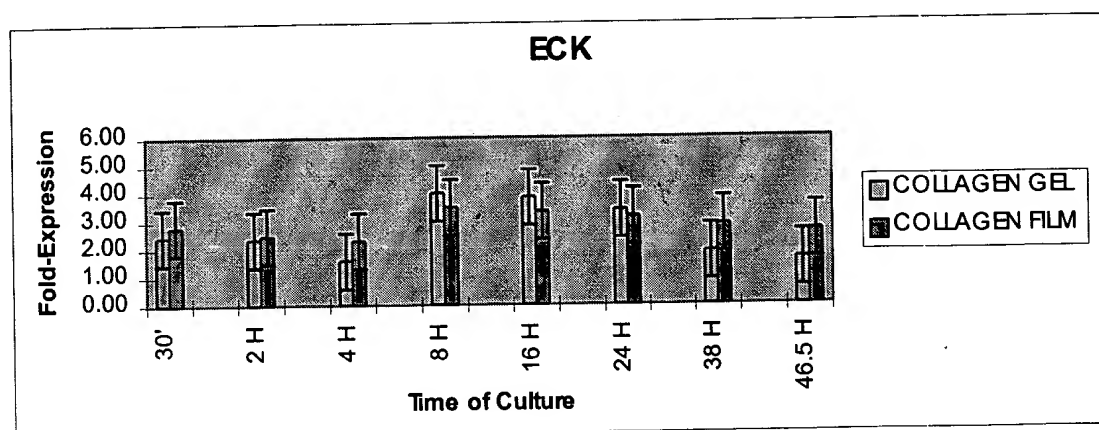


FIGURE 17: PA17 (OX-40)

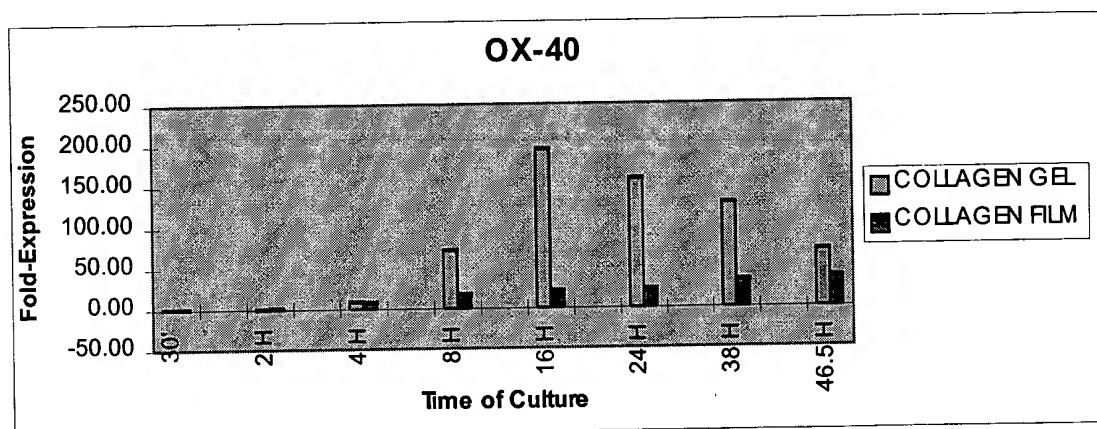


FIGURE 18: PA18 (gp130)

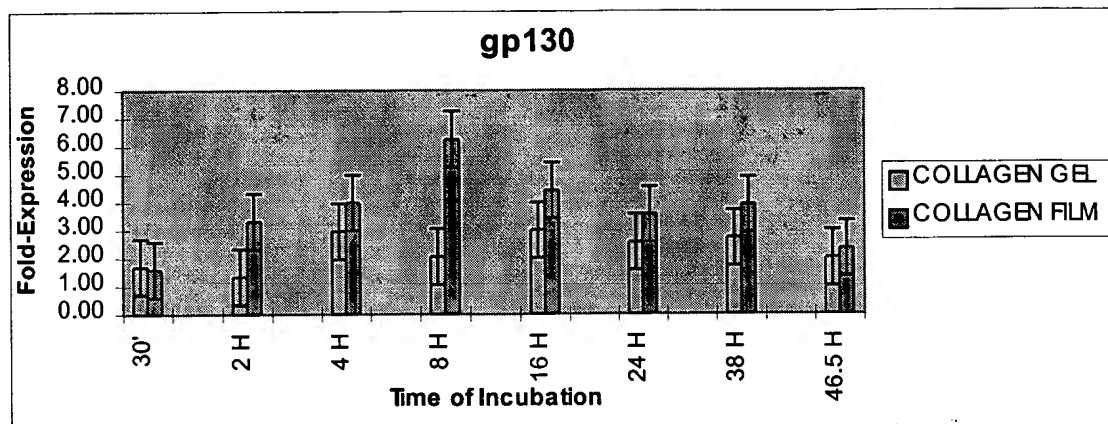


FIGURE 19: PA19 (CD82)

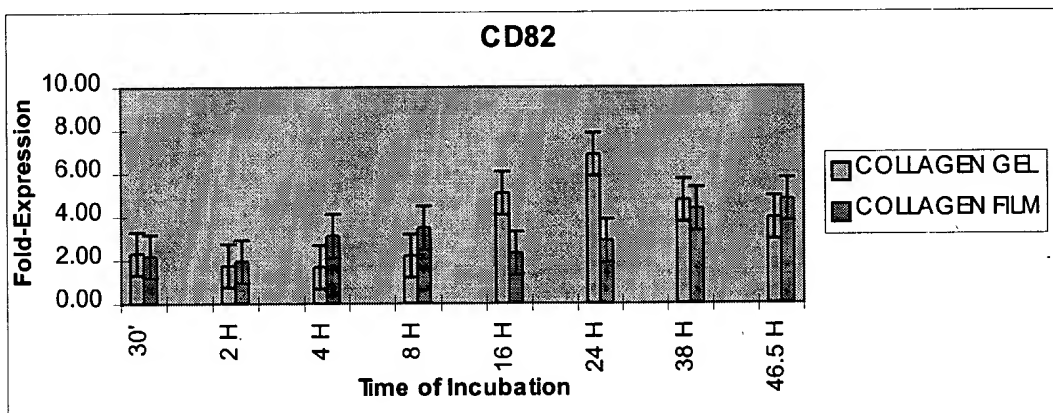


FIGURE 20: PA20 (PRZ)

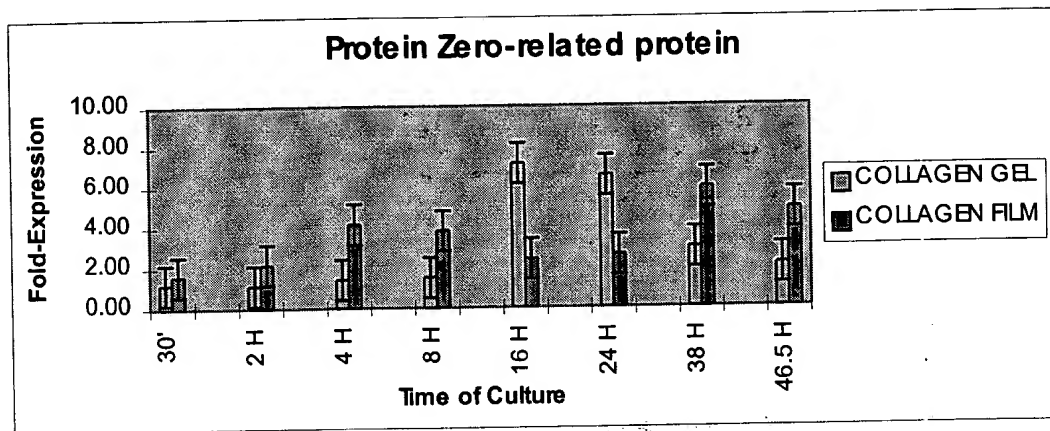
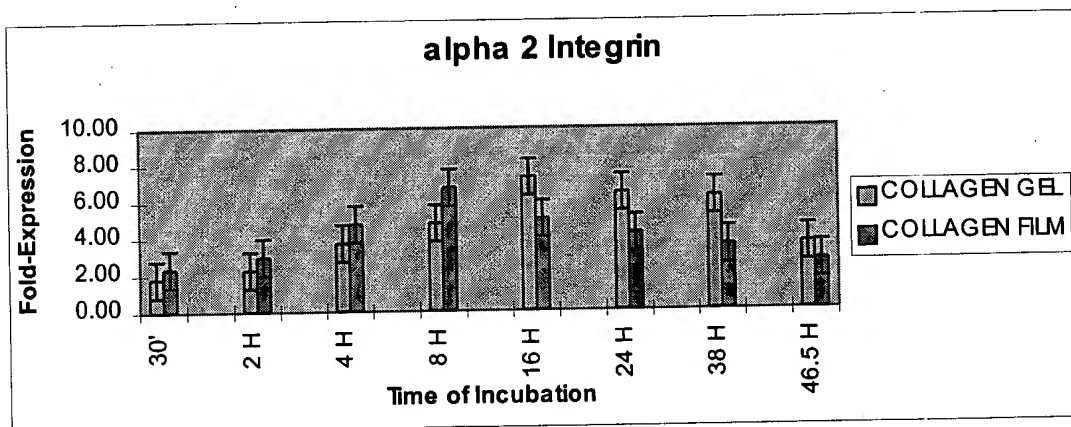


FIGURE 21: PA21 (alpha 2 integrin)



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FIGURE 22: PA22 (PlGF)

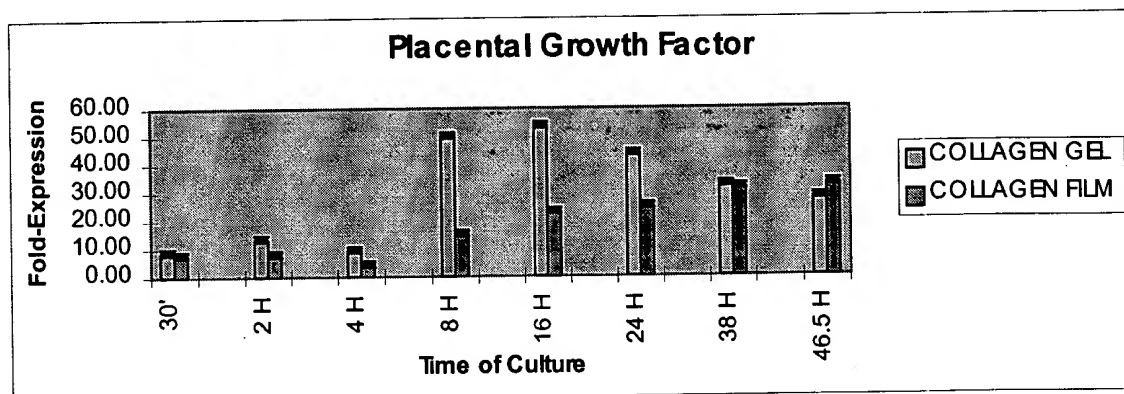
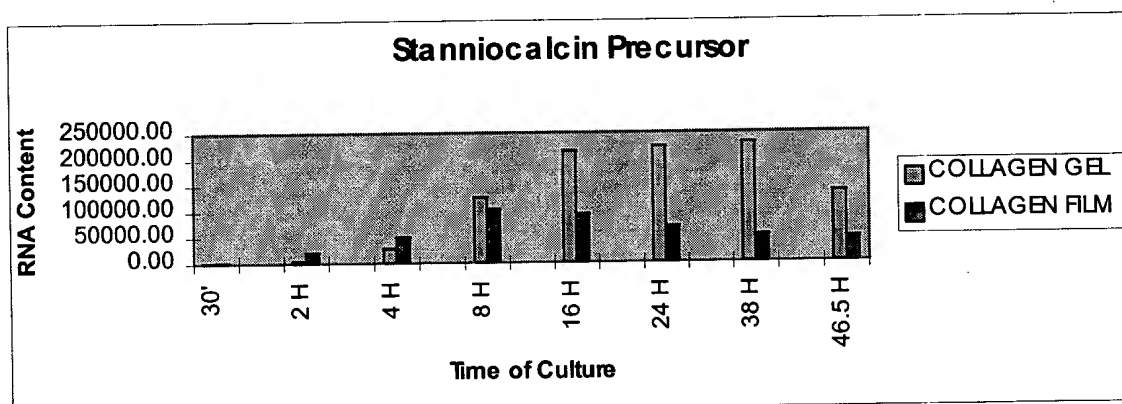


FIGURE 23: PA23 (stanniocalcin precursor)



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FIGURE 24: PA24 (FGF-16)

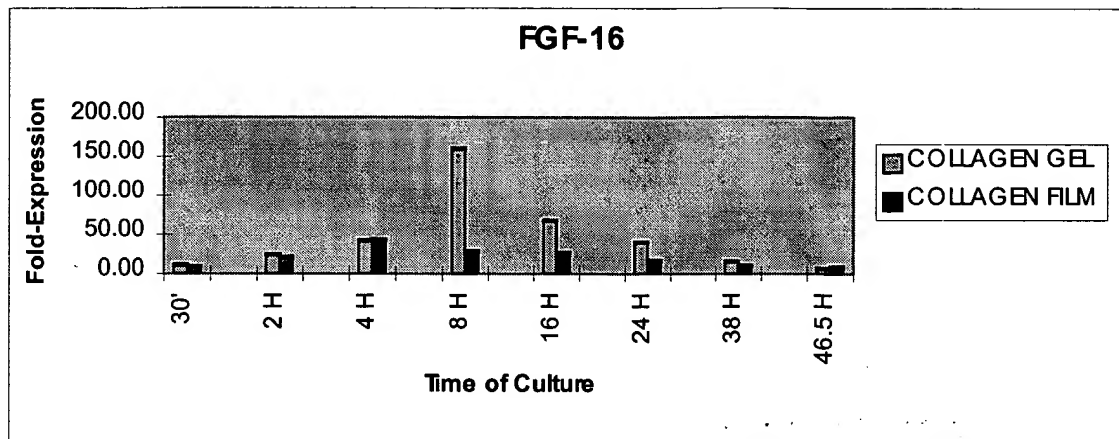


FIGURE 25: PA25 (White Protein Homolog)

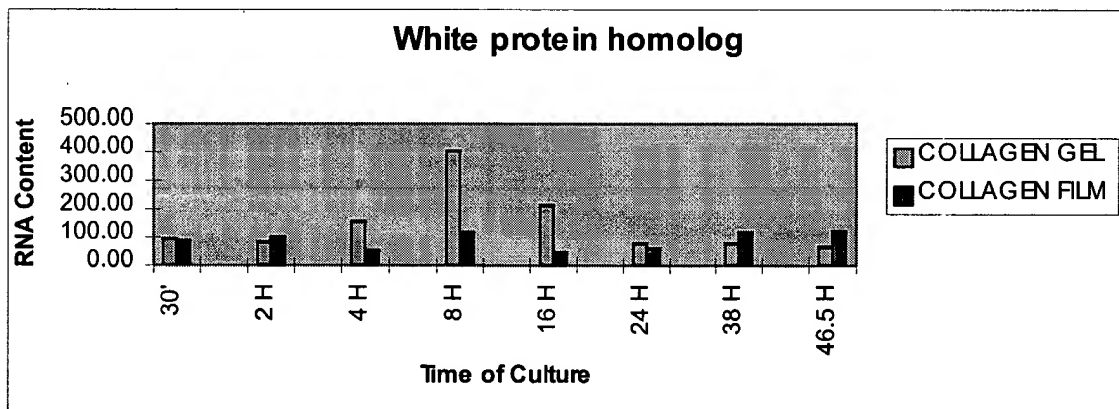
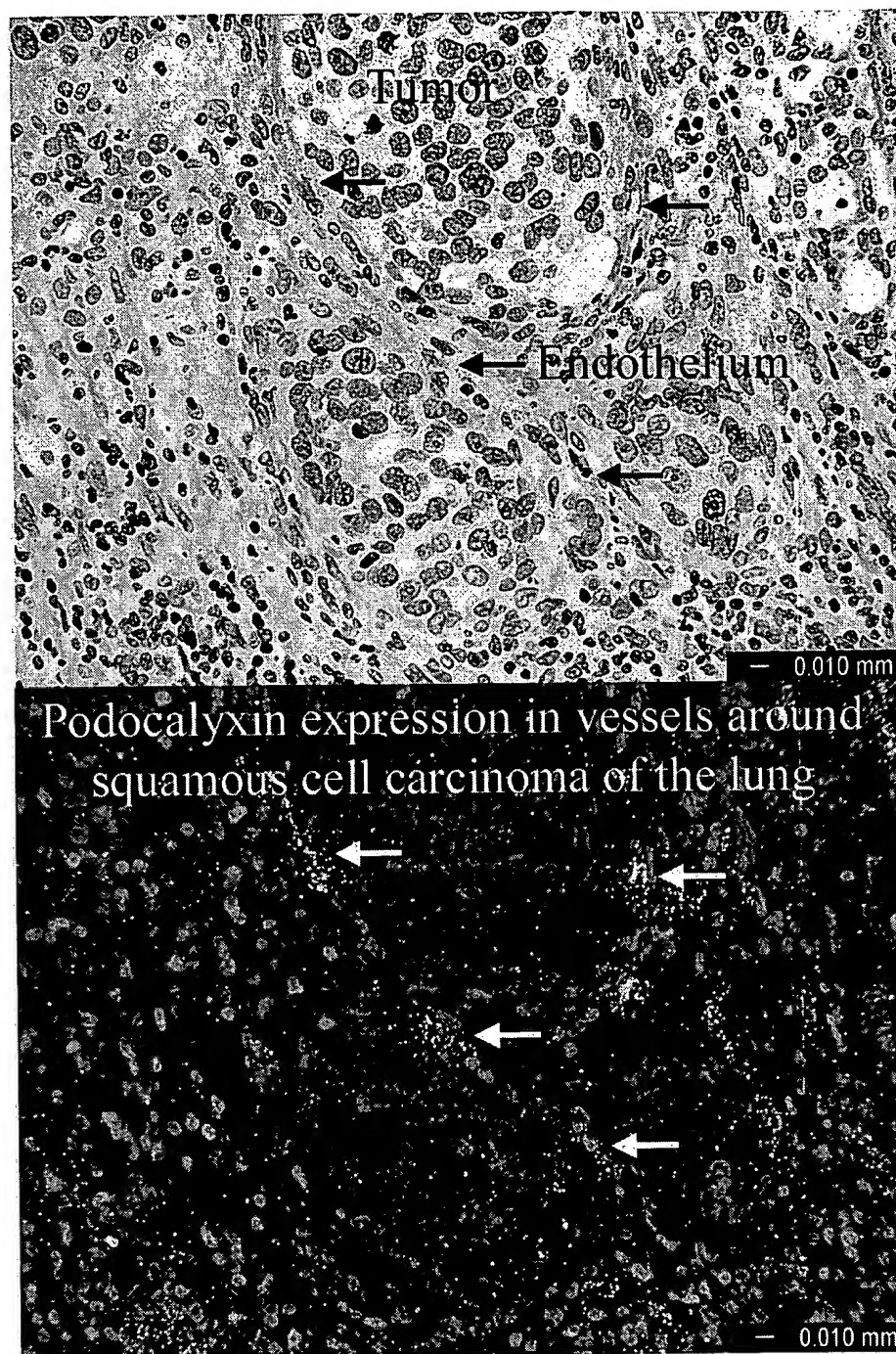


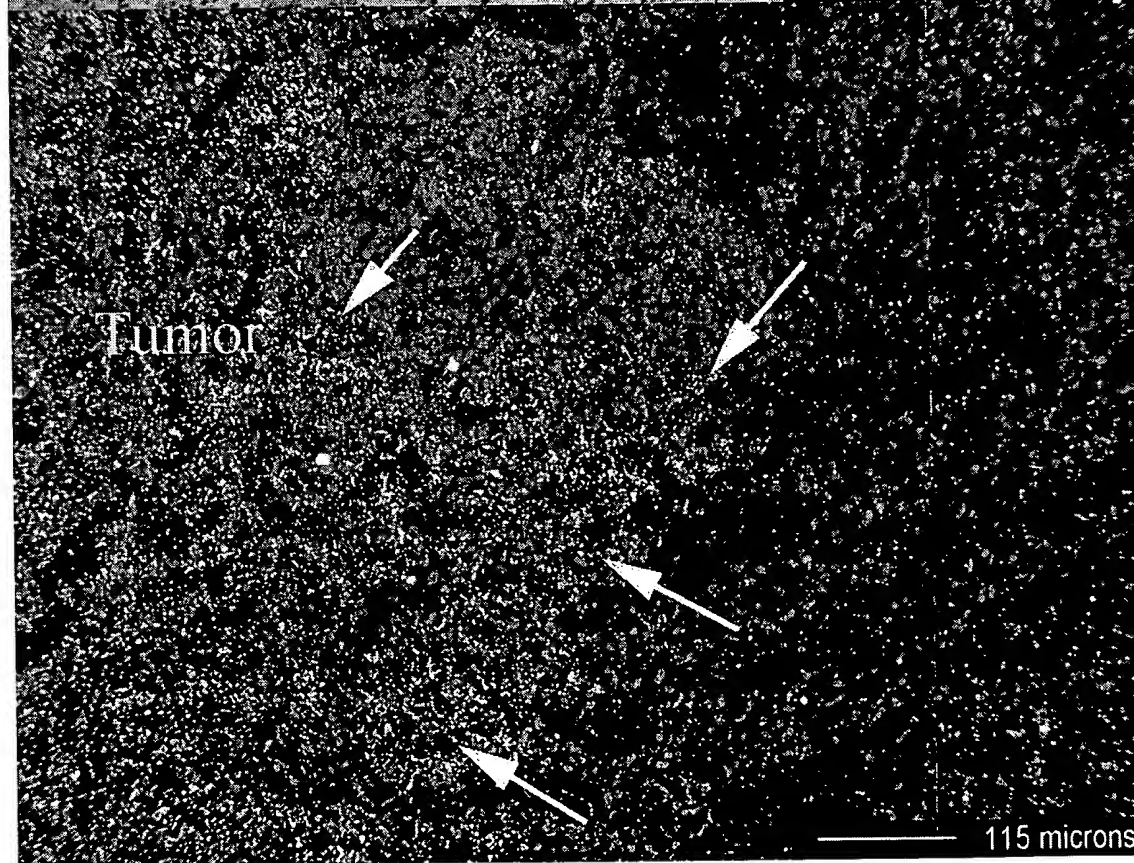
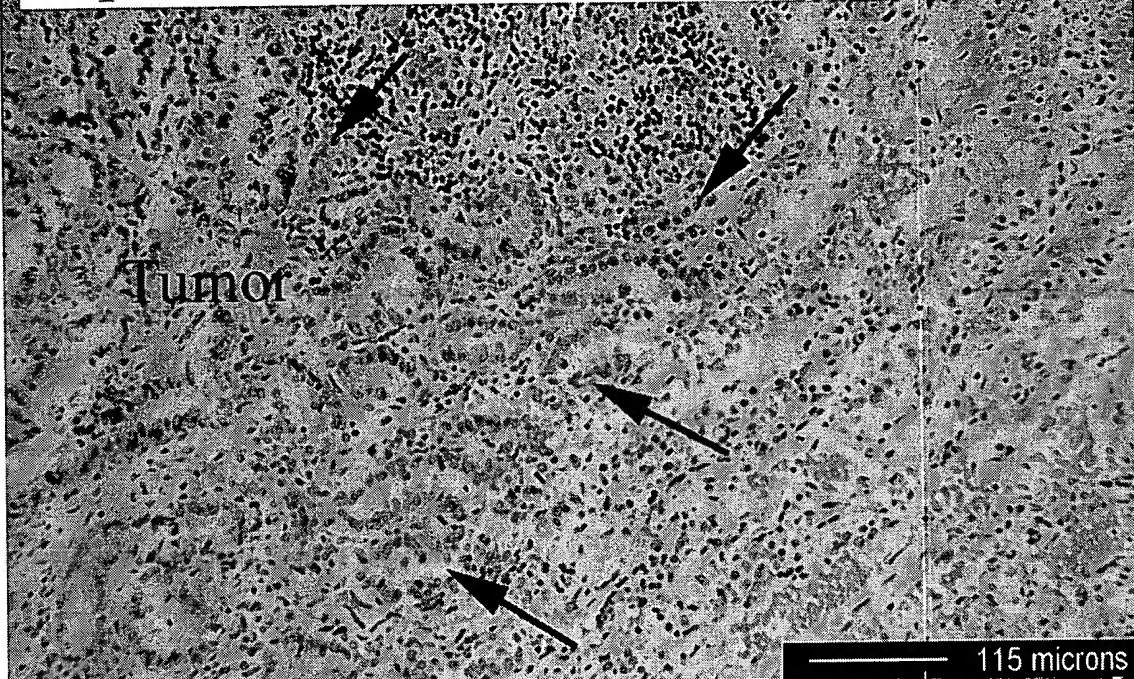
FIGURE 26



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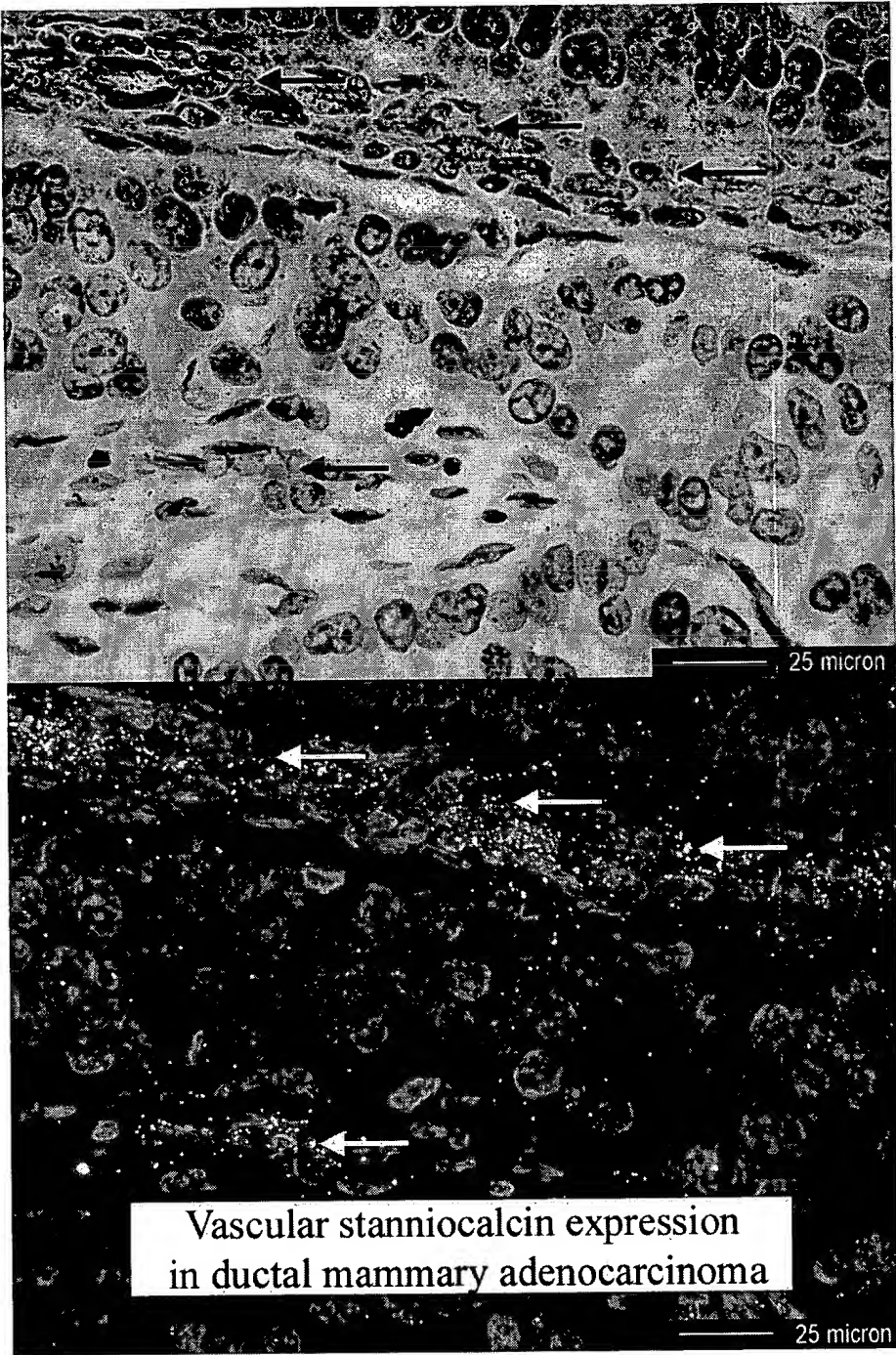
FIGURE 27.

DNA 76510 (Protein Zero Homolog) Expression in Pulmonary Adenocarcinoma



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FIGURE 28



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FIGURE 29

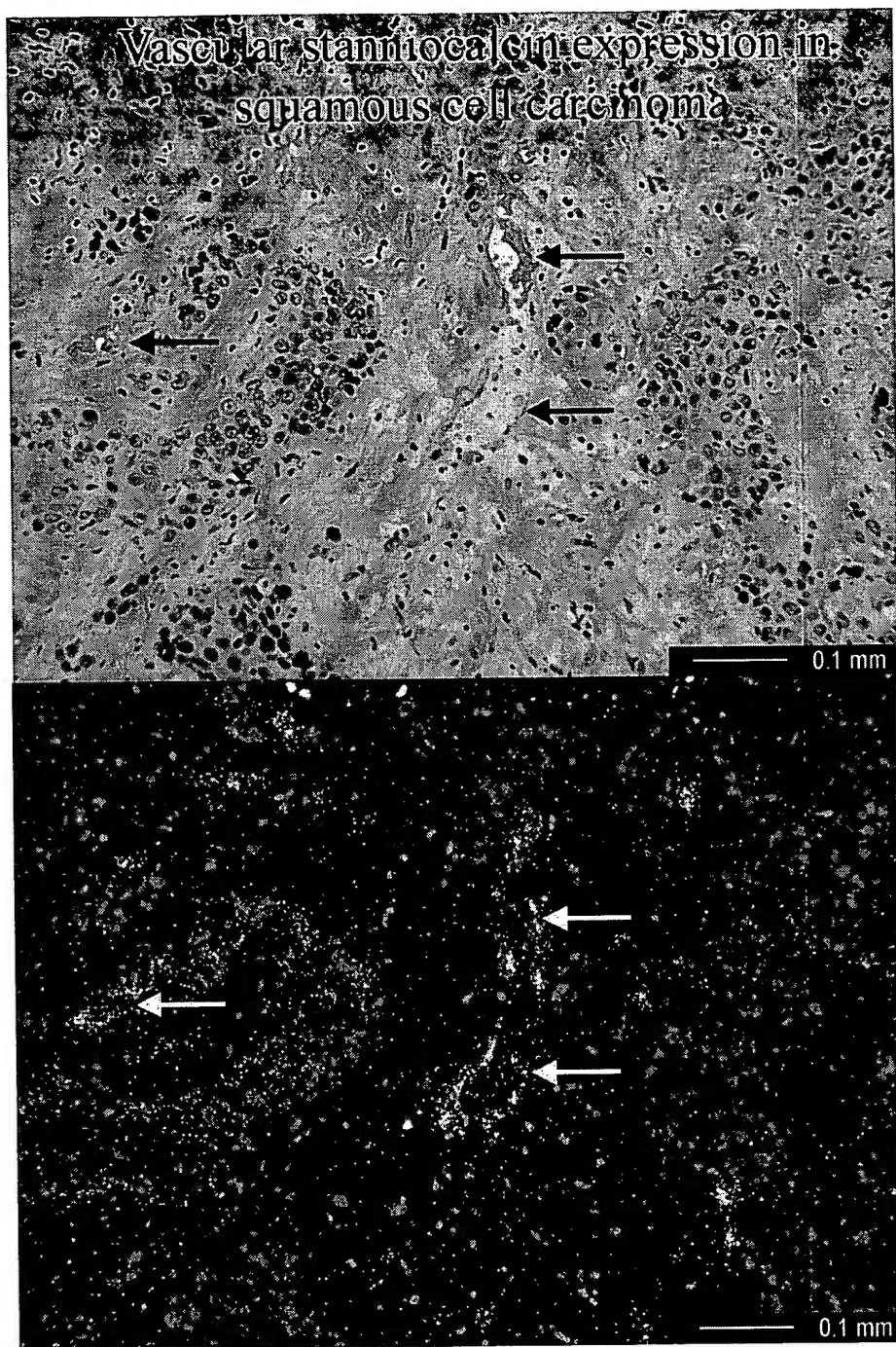
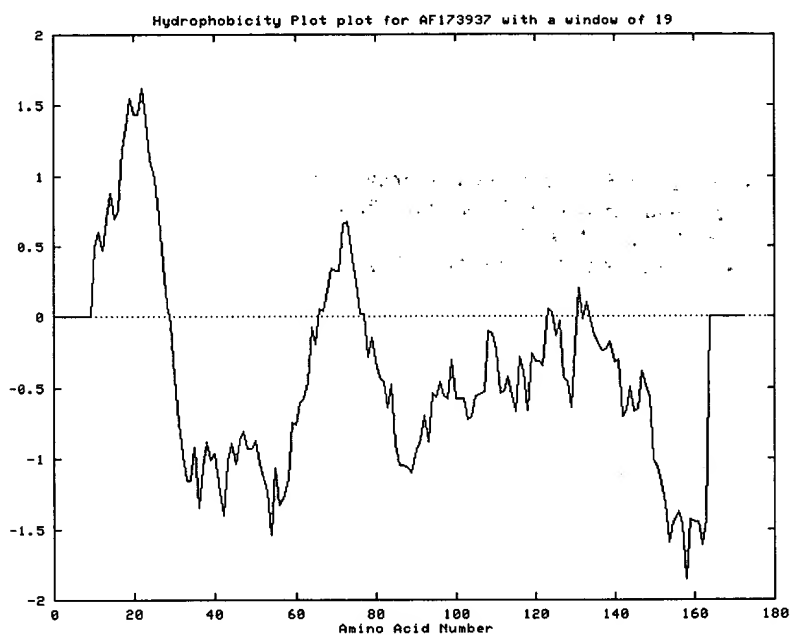


FIGURE 30

1 HVGPAPRRRLRPLAALALVLALAPGLPTARAGQTPRPAEROPPVRLFTTEELARYGGELEDQPIYLAVKGVVFDVTSCKE
01 FYGRGAPYNALTGRDSTRGVAKHSLDPADLTHDTTGLTAKELEALDEVFTKVYKAKYPIVGYTARRILNEDGSPNLDPKP
161 EDQPHFDIKDEF (SEQ. ID. NO.: 72)

FIGURE 31



0070350-103100

FIGURE 32

Sequence type explicitly set to Protein
Sequence format is Pearson
Sequence 1: 55193557_EXT 153 aa
Sequence 2: 11725372_EXT 161 aa
Sequence 3: AF173937 172 aa
Start of Pairwise alignments
Aligning...
Sequences (1:2) Aligned. Score: 83
Sequences (1:3) Aligned. Score: 83
Sequences (2:3) Aligned. Score: 90
Start of Multiple Alignment
There are 2 groups
Aligning...
Group 1: Sequences: 2 Score:1860
Group 2: Sequences: 3 Score:1948
Alignment Score 2458
CLUSTAL-Alignment file created [/data4/genetools/lrastelli5501clustalw]

Multiple Alignment:

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11725372_EXT  -----RPLAALALALALVR-VFSARAGCMPRPAERGPPVRLFTEELARYSGEEE
AF173937      MVGPAPRRRLRPLAALALVLAAPGLFARAGCTPRPAERGPPVRLFTEELARYGEEEE

55193557_EXT  DQPIYLAVKGVVFDVTSKGKEYGRGAPYNALAGKDSSRGVAKMSLDPADLTHDTTGLTAK
11725372_EXT  DQPIYLAVKGVVFDVTSKGKEYGRGAPYNALAGKDSSRGVAKMSLDPADLTHDI-SGLTAK
AF173937      DQPIYLAVKGVVFDVTSKGKEYGRGAPYNALTGKDSIRGVAKMSLDPADLTHDTTGLTAK
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